



OIPE

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## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/887,272A

DATE: 03/26/2002 P.S  
TIME: 14:20:44

Input Set : D:\Sequence.txt

Output Set: N:\CRF3\03262002\I887272A.raw

4 <110> APPLICANT: Hou, Yu-Ming  
5 Quan, Sheng  
6 Chang, Hur-Song  
7 Zhu, Tong  
8 Whitham, Steve  
9 Goff, Steve  
10 Glazebrook, Jane  
11 Chen, Wenquiong  
12 Katagiri, Fumiaki  
13 Xie, Zhiyi  
14 Tao, Yi  
15 Zou, Guangzhou  
16 Cooper, Bret  
18 <120> TITLE OF INVENTION: PLANT GENES INVOLVED IN DEFENSE AGAINST  
19 PATHOGENS  
21 <130> FILE REFERENCE: 1360.003US2  
23 <140> CURRENT APPLICATION NUMBER: US 09/887,272A  
24 <141> CURRENT FILING DATE: 2001-06-23  
26 <150> PRIOR APPLICATION NUMBER: 60/213,634  
27 <151> PRIOR FILING DATE: 2000-06-23  
29 <150> PRIOR APPLICATION NUMBER: 60/214,926  
30 <151> PRIOR FILING DATE: 2000-06-23  
32 <150> PRIOR APPLICATION NUMBER: 60/261,320  
33 <151> PRIOR FILING DATE: 2001-01-12  
35 <150> PRIOR APPLICATION NUMBER: 60/264,353  
36 <151> PRIOR FILING DATE: 2001-01-26  
38 <150> PRIOR APPLICATION NUMBER: 60/273,879  
39 <151> PRIOR FILING DATE: 2001-03-07  
41 <150> PRIOR APPLICATION NUMBER: 09/887,271  
42 <151> PRIOR FILING DATE: 2001-06-22  
44 <160> NUMBER OF SEQ ID NOS: 6813  
46 <170> SOFTWARE: FastSEQ for Windows Version 4.0  
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49 <211> LENGTH: 636  
50 <212> TYPE: DNA  
51 <213> ORGANISM: Arabidopsis thaliana  
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56 tcagaaaagg ccactgaaca agatacagta tcatcaattg gggtaaagaa acctccagt 180  
57 gactctccag ccactactaa tgctgctagc ggtcgacttg tttatgtccg gagaagagt 240  
58 gaggttgaca catccaaagc cgctgctagc actactaatc cgaatccgcc tcctactaag 300  
59 gcaccacccc agataccttc atcacccgca caggcacagg cacaggaacc aaccctacc 360

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60 agccataagt tagactggga agagcggttac cttcatcttc aaatgcttct taacaagctt 420
61 aatcaatctg atcggaccga tcatgttcag aatatgtttc ctctgttatt agtgctttgg 480
62 tcactttcct ccgccagct tagcaagcat gctgttgact tggaaaagag gtctattcag 540
63 ttctctctcg aggaagcgag agagatgcag cgcgtagcag ctttaaactg gttaggaag 600
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67 <211> LENGTH: 1400
68 <212> TYPE: DNA
69 <213> ORGANISM: Arabidopsis thaliana
71 <400> SEQUENCE: 2
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74 aacaaacgat ccaagctatg gagcaatgga acaagtgtct ctaagtttag tctacttgaa 180
75 gagccaagtc ctttgggggt gagtttaaaa aagagcccat cttttcaaga actgattgag 240
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77 ggtggtgtag gaaccgttga gaagcttaaa gtttctaact ttctgctac tattctcagg 360
78 attggtcaat gggagtataa gtcaaggat gaaggtgatt tgggtggcga atgttacttt 420
79 gcaaaacata aacttggtg ggaagtgtt gaacaaggtc ttaagagcaa gattgagatt 480
80 cagtggctcg atattatggc tttgaaggct aatttgccag aggatgaacc tggaaccctg 540
81 actattgtgc tggctaggcg gccattgttt ttcagagaaa ctaatccgca gcctagaaaa 600
82 catactttgt ggcaggcgac atcggatttt actgatggtc aagctagcat gaacaggcaa 660
83 cttttctgc agtgtcccc agggattatg aacaacatt ttgagaagct tgttcaatgt 720
84 gatcatcgct tgttctgtct aagccggcag ccagagataa acttgccgc accgttcttt 780
85 gattcacgac tatctatatt tgaggatccc tcagtgtctg gatctcataa tattgcatct 840
86 cctgttgggg ctcatcatc atcagaacat gtgtctctat ctcatgacgc actatcgctt 900
87 agctcagtga tggatgctcg tgcaatcgaa ggagttggtg gttccattga ttcgaggaat 960
88 acaaacgggt ggagtcagat aaaaatgcct ggactacacc aatctatctc gatgaacgat 1020
89 ttctttacgt ttttgtcaga tcaagcttgt gaaaacaacc aagagtttga ggaaatgaaa 1080
90 cagttaactgc taagtgacaa cactcaaacc gatccatctg atgagaagtc tgtcatgtca 1140
91 aaggtgaatt ctttctgcaa cctcttgcaa tctgctgcaa actcacagct caatatcgaa 1200
92 actgctgata ccgaaagagt tgttggggtc gataacaata gacacatgcc cgaaggaggc 1260
93 aaaagagttg ttgacctgc ttcaagcagc aagccactac aagggatgtc aaggaaagac 1320
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95 aacatttcag aagaagacta 1400
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98 <211> LENGTH: 1710
99 <212> TYPE: DNA
100 <213> ORGANISM: Arabidopsis thaliana
102 <400> SEQUENCE: 3
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104 gcttacgtca cgacggcgaa ggcggtgagc gttccggttc aagtctctcc ggcgactgtg 120
105 gtttccgaag gattaggtaa agatgcgttg atctcatggt tcagaggaga attcgctgcg 180
106 gcaaacgcga taatcgacgc gatgtgctcg catctaagga tagctgaaga agcggtatct 240
107 ggatcggagt acgaagctgt attcgcggcg atccatcgcc gtcgtctgaa ttggatccct 300
108 gttcttcaga tgcagaagta tcattccatc gctgaagtag cgatcgagct tcagaaagtg 360
109 gcggtctaaga aagcggagga tctgaaacag aagaaaacag aggaggaggc ggaggaagat 420
110 ctgaaagagg tgggtggcgac ggaggaagag gaagtgaaga aagagtgtt taacggcgag 480
111 aaagtaacgg agaatgacgt aaacggagat gtagaagacg ttgaagatga ttcaccaact 540
112 agtgatatca ctgattcagg ttctcatcag gatgttcacc aaactgttgt tgcggatact 600

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Input Set : D:\Sequence.txt

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113 gcacatcaga ttatatgcca tagccatgaa gatttgtgatg cacgttcatg tgagatcaaa 660
114 cctatcaaag gtttccaagc taaagaacaa gtcaaaggcc acactgtgaa tgttgtgaaa 720
115 ggactaaagc tgtatgaaga gttacttaaa gaagatgaga tatctaaatt gttagacttt 780
116 gtggcagaac tacgagaagc tggcataaac ggggaagcttg caggtgagag ctttatactg 840
117 ttcaacaaac agattaaagg gaacaagaga gagctgattc agcttggtgt ccccatcttt 900
118 ggatcatgtca aggcggatga aaactctaac gacaccaaca actctgttaa catcgagcca 960
119 attccaccac ttcttgagag tgtcattgat cactttgtca catggagact catcccagaa 1020
120 tacaagagac ctaacggctg tgtcatcaac ttctttgaag aggggtgaata ctacagcct 1080
121 ttctcacaac cacctcaactt agaacaacca atctccactc ttgtcctctc tgaatcaaca 1140
122 atggcctatg gacgcattct ctcaagtgaac aacgaaggca acttcagagg acctttgaca 1200
123 ctctctctca aacaaggatc ttgttggtg atgagagggga acagtgcaga catggcaaga 1260
124 catgtaatgt gtccatcaca aaacaaaaga gtaagcatca cattctttcg tattcgacct 1320
125 gacacatata ataaccattc acaaccaaac agtctcgcga acgacggtgt catgacaatg 1380
126 tggcaaccct accaaatgac accaactcca ttctcaatg gttatgatca ttcaattgac 1440
127 atgatgccaa aacttggagt cctacgtcct ccaatggtca tgatggcacc accaccggtt 1500
128 caaccaatga tattaccaag tcccaatgtg atgggaaccg gtggtggtac cgggtgttttc 1560
129 ttaccatggg cctctgttaa tagctcaaga aaacatgtga agcatttgcc tccacgcgcg 1620
130 cagaagaagc gattacttcc gcttctcctc gctgcttctt cttctccagc tggaggatcc 1680
131 acctctgagc ctgtgatcac tgtaggttaa 1710

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133 &lt;210&gt; SEQ ID NO: 4

134 &lt;211&gt; LENGTH: 462

135 &lt;212&gt; TYPE: DNA

136 &lt;213&gt; ORGANISM: Arabidopsis thaliana

138 &lt;400&gt; SEQUENCE: 4

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139 atgggtgttc ttgatcacgt ctctgaatat ttogattgtt ctcatggaag ctccaagaga 60
140 cacaaaagtc tacagacggt ggatgtgagg gttttgatag attgtgaagg atgcgagagg 120
141 aaagtaagga gagctttaga aggaatgaga ggaataagag atgtaaccat cgagccaaat 180
142 gctcagaaag tgacagtggg tgggtacgtt gaaccaaaaca aagtgggtggc tcgtatcatt 240
143 caccgaaccg gtaaaagagc agagctatat ctttctgttc cttacgacgt tgtggctcat 300
144 ccttacgcgt ctgggtgttta cgataacaga gccccgactg ggtacgttag gaacaccgag 360
145 tatgatccac atgtgtcacg tctcgcacgt gctagctcca ctgaggttcg ttatactacg 420
146 gcgttttagc acgagaacgc ctccgcttgt gttgttatgt ga 462

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148 &lt;210&gt; SEQ ID NO: 5

149 &lt;211&gt; LENGTH: 885

150 &lt;212&gt; TYPE: DNA

151 &lt;213&gt; ORGANISM: Arabidopsis thaliana

153 &lt;400&gt; SEQUENCE: 5

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154 atgttcagaa gaaggaatgc tcaccaactc gacaatgatg attctcaaca agataacaag 60
155 gttagagaat tgaaatccgc gataggacca ttatcgggac atagttagt tttctgctct 120
156 gatgcttctc tgaggagatt tctagatgct cgtaattggg atgtggaaaa agccaagaaa 180
157 atgatccaag agactcttaa atggagatca acttataaac ctcaagagat ccgttggaat 240
158 caagtagcac atgaagggtga gaccggaaaa gtttcaagag ctagttttca tgatcgacaa 300
159 ggtagagtag tgcttataat gagaccagcc atgcagaact caacatcaca agaaggtaat 360
160 atcaggcatt tgggtgtatct tcttgaaaat gcaatcataa atcttcccaa gggacaaaaa 420
161 caaatgtctt ggctcattga tttcactggg tgggtctatg ctgttaatcc tcctatgaaa 480
162 acaacacgcg aaatcattca cattctacag aactattacc ctgagagact cggtatcgcc 540
163 tttctctaca atccaccaag actcttccaa gcagtctaca gggctgctaa gtacttcttg 600
164 gaccacgta cagctgaaaa ggtgaaattt gtgtacccaa aagacaaagc aagtgatgaa 660
165 ctgatgacga cacattttga cgttgagaat ctcccaagg aattcggagg tgaagcaaca 720

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166 ctagagtatg atcatgaaga tttctcacga caaatgtatg aagacgatct caaaaccgca      780
167 aaatactggg gactagaggg gaagcactat ccgaaaacaa acggtttctc tccatccgat      840
168 gttgttcctg agccggcgat agaaatcgca tcagcagcta gctga                      885

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170 &lt;210&gt; SEQ ID NO: 6

171 &lt;211&gt; LENGTH: 1224

172 &lt;212&gt; TYPE: DNA

173 &lt;213&gt; ORGANISM: Arabidopsis thaliana

175 &lt;400&gt; SEQUENCE: 6

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176 atggaagtgt cgaaaaagaa aggtggagat tttcagcaat gtcacgaatt gattcccggg      60
177 ttaccttccg aactagcttt ggagtgtttg gtcagagttc cgttccaatt ccaatccgcc      120
178 atgagatccg tttgcogttc ttggcgtagc ttactctccg actcttcttt tatccaagaa      180
179 cggcggagat gcggcaaaac agagcttctc ctctgcctcg ttcaaccgct aacgccgcca      240
180 attccagcgt ctaaatcggt tgacgagaca ttaatggttg acgagaaaaa atcagaggat      300
181 gagtcacacc cgcgcgtttt ctgcacgcgc cgttttggat tgagcgttta caacgctgcg      360
182 atgtccacgt ggcacgcgtg tgcgtttcca gaggaggagc agatcccgtt tttctgcgag      420
183 tgcgtcgtgc ttcaggacgc cggaagatt ctgctcatcg gcggttgga tccggagacg      480
184 ttacagccga cgagagacgt ttacgtttct gaattcgccg gaaggaagtg gagacgaggt      540
185 gcgccaatga aggaatcacg atcattcttc gcctgcgcct ccgtaagtcc aacgaaggta      600
186 tacgtcgccg gaggtcacga cgatcagaaa aacgctttac gctcggcgga ggtgtacgac      660
187 gtcgagaaaag acgagtggtc gtcggttact ccgatgacgg aaggaagaga cgaatgtcaa      720
188 ggattcgccg ttggtatggg tctaaggttt tgcgtgctga gcggttacgg aacggaatct      780
189 caaggaagat tccgacttga cggagaaatt tacgatccag cgacagattc atggtcgaga      840
190 atcgacaacg tctggcgatt tccagatact agcccagag gtcgcaccgc cggagacttt      900
191 agaagctcct ctacgttatg gtgtttcaca gacactgatt tacagagtga acgtcgatgg      960
192 gaaactaacg atgattcgag aaatttgaaa ctggatcttc aatctataca gcttccgatg     1020
193 acaggaagct cagttttcgc cggaagctta ggaggtgaat cgggtggaat gattggcggg     1080
194 aaaagagaga gtgaaggtga aggtgaagga ggagtgatga tgaagatgac gactgagaag     1140
195 aaaatgggaa aatggagtca tcatgttcat ataccttggt atttctctac tcttccattt     1200
196 tcgcatgctt caatctatgt ttga                                           1224

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198 &lt;210&gt; SEQ ID NO: 7

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201 &lt;213&gt; ORGANISM: Arabidopsis thaliana

203 &lt;400&gt; SEQUENCE: 7

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205 tctccgtcgc ttcagtcatt tctccggcg gcggcaatcc gttcttctcc atcaccgatc      120
206 ttcagaaaag ctccagcggg gttcaacaac ggcgacgaat gtctctctcc cggcggcgtc      180
207 tgcaatccgt cgttgggtcca cgtggcgatc acgttagacg tagagtacct gcgtggctca      240
208 atcgacgcgc ttaactcgat ccttcagcac tcggtgtgtc cagagagcgt cttcttcac      300
209 ttcacgcgcg tctccgagga aacaaacctg ttggagtgcg tggtagatc ggttttcccg      360
210 agactgaaat tcaatattta cgattttgcc cctgagacag ttcgtggttt gatttcttct      420
211 tccgtgagac aagctctcga gcagcctctg aactacgcta gaagctactt agcggatctg      480
212 ctggagcctt gtgttaaccg tgtcatatac ttggattcgg atcttgtcgt cgtcgatgac      540
213 atcgctaagc tttggaaaac tagcctaggc tcgaggataa tcggagctcc ggagtattgt      600
214 cacgcgaatt tcacgaaata cttcaccgga ggattctggt cggaggagag attctccggg      660
215 accttttagag ggaggaagcc atgttacttc aacacagggt tgatgggtgat agatcttaag      720
216 aaatggagaa gaggtggtta cacgaaacgt atcgagaaat ggatggagat tcagagaaga      780
217 gagaggattt acgaactagg ctgcgttcca ccgtttcttc tagttttctc cggtcacgtg      840
218 gctcccatct ctcaccgggt gaaccagcat ggacttggtg gtgacaatgt tagaggtagc      900

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221 ttgtatcgac attcgcattg                                     1040
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224 <211> LENGTH: 765
225 <212> TYPE: DNA
226 <213> ORGANISM: Arabidopsis thaliana
228 <400> SEQUENCE: 8
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231 cttaaagagc aattctttgt tcattgcaag tttcatggag actctcataa gagtgaatgc      180
232 aatatgtatt gcttagactg caccaatggc ccactttgct ctctttgtct tgctcatcac      240
233 aaggatcatc gcaccattca gataaggaga tcttcttata atgatgttat aagggtgaat      300
234 gagatacaaa agtatcttga tataggtggg atccaaacct atgtgatcaa tagtgctaaa      360
235 gtcgtcttcc tgaatgagag gcctcagcct aggcctggga aaggagtgaac caatacctgc      420
236 aaggtttgct atcgtagcct cgttgatgat agcttccgct tctgctctct tggctgcaag      480
237 ttccacctaa cttctccttg catttcagta aactcttttc tgatacttac tttctctatg      540
238 tttttgggaa atgaaatcaa gattgctgga acatcaagag gctttgaaaa gggaaggag      600
239 aacctgctaa tggaacaga ggattcaagt agcagcattg caattgggaa gaacattaca      660
240 aatctccaga gtttcagccc atcaaacacct ccacttacta catctagtaa ctgcagaatc      720
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244 <211> LENGTH: 1398
245 <212> TYPE: DNA
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251 cccgttaacc cgattcagaa ccggtttaac tcttcccaag ctcccaatcc gactcctaga      180
252 cccaaacca atccaaatcc gttgcctgag aaacccttat cctcctccga tcagaagatt      240
253 agtgggtcaa cgagaaaccc ggatcatgac ccggttcgtg ctccacaaga tgggtttgac      300
254 ccgtacggtg ctttcataac gtcttcgaac agatctcaga acggatattc tctgagcatg      360
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258 agctctattg atgttttgtc taggttgttt aagaacatag tgaaagaacc tgagaatgct      600
259 aagtttagaa aagttagaat gagtaatgct aagattaaag aagctattgg tgatgttgct      660
260 ggaggtgtcg agcttcttga gttggttggg tttgaattga aagaggaaaa tgatgagatt      720
261 tgggctgtta tggatgttcc tagtgaagaa caatctatat tgatcaataa agttgtcggg      780
262 tatctagaaa agagaagac agagagttca ggaatctcag ctacaggttat ggaacctgtg      840
263 gctccaaaga agattgatag agagaatggg tgtttcttgg ttcagattcg tgtcttttct      900
264 tccggtttctg aaaacgtagc atcgagaata gaggtacctg attcgttcta tagtctctcg      960
265 gcggatgaaa taaagagaga agcggattta cggaggaaaa agattgcgga gtcacagctt      1020
266 ttgattccaa gatcttataa ggagaaacaa gcgaaagcag ctagaaagag atacaaaaga      1080
267 agtatgataa gagtgcagtt tcttgacgga gttgtgcttc aagggtgtgtt tgctccatgg      1140
268 gaacctactt ttgctctcta tgagtttgtg agctctgctt tgaaagaacc aagtttgcag      1200
269 tttgagcttc tggatcctgt gctggttaaa cggcgtgtga ttccacatac tccagccccg      1260
270 ggacagaaac cgataacggt agaggacgaa gagcttgctc cgtcagcgct catcagggtc      1320
271 agaccgatcg aaacagactc tctcgtcttt acaggactcc gcaatgaact cctggagatc      1380

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Use of n and / or Xaa has been detected in the Sequence Listing. Review the Sequence Listing to ensure a corresponding explanation is present in the <220> to <223> fields of each sequence using n or Xaa.

## VERIFICATION SUMMARY

DATE: 03/26/2002

PATENT APPLICATION: US/09/887,272A

TIME: 14:20:45

Input Set : D:\Sequence.txt

Output Set: N:\CRF3\03262002\I887272A.raw

L:2054 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (69) SEQUENCE:  
L:2559 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (87) SEQUENCE:  
L:2800 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (96) SEQUENCE:  
L:3031 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (105) SEQUENCE:  
L:4197 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (146) SEQUENCE:  
L:4202 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (147) SEQUENCE:  
L:4234 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (149) SEQUENCE:  
L:4348 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (155) SEQUENCE:  
L:4378 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (157) SEQUENCE:  
L:4481 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (161) SEQUENCE:  
L:4615 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (167) SEQUENCE:  
L:6959 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (245) SEQUENCE:  
L:7632 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (268) SEQUENCE:  
L:7637 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (269) SEQUENCE:  
L:7875 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (276) SEQUENCE:  
L:7880 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (277) SEQUENCE:  
L:7885 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (278) SEQUENCE:  
L:8251 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (293) SEQUENCE:  
L:8711 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (311) SEQUENCE:  
L:8944 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (320) SEQUENCE:  
L:8980 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (322) SEQUENCE:  
L:9256 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (332) SEQUENCE:  
L:9299 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (335) SEQUENCE:  
L:9629 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (349) SEQUENCE:  
L:9881 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (361) SEQUENCE:  
L:10052 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (368) SEQUENCE:  
L:10491 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (379) SEQUENCE:  
L:10828 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (396) SEQUENCE:  
L:10911 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (400) SEQUENCE:  
L:11686 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (430) SEQUENCE:  
L:11924 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (442) SEQUENCE:  
L:11970 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (445) SEQUENCE:  
L:12200 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (455) SEQUENCE:  
L:12205 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (456) SEQUENCE:  
L:12257 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (459) SEQUENCE:  
L:12410 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (466) SEQUENCE:  
L:12477 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (469) SEQUENCE:  
L:12666 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (477) SEQUENCE:  
L:12671 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (478) SEQUENCE:  
L:13966 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (511) SEQUENCE:  
L:14007 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (514) SEQUENCE:  
L:14136 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (519) SEQUENCE:  
L:14266 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (524) SEQUENCE:  
L:14363 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (527) SEQUENCE:  
L:14407 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (529) SEQUENCE:  
L:14510 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (534) SEQUENCE:  
L:14595 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (538) SEQUENCE:  
L:14852 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (547) SEQUENCE:

## VERIFICATION SUMMARY

DATE: 03/26/2002

PATENT APPLICATION: US/09/887,272A

TIME: 14:20:45

Input Set : D:\Sequence.txt

Output Set: N:\CRF3\03262002\I887272A.raw

L:15097 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (555) SEQUENCE:  
L:15160 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (558) SEQUENCE:  
L:30567 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1955  
L:30568 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1955  
L:30570 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1955  
L:30596 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1957  
L:30623 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1959  
L:30638 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1960  
L:30656 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1961  
L:30696 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1964  
L:30718 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1965  
L:33868 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2003  
L:33892 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2004  
L:33917 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2005  
L:33942 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2006  
L:33946 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2006  
L:33951 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2006  
L:33952 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2006  
L:33954 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2006  
L:33956 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2006  
L:33957 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2006  
L:33958 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2006  
L:33983 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2007  
L:33984 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2007  
L:34007 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2008  
L:34047 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2010  
L:34049 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2010  
L:34050 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2010  
L:34109 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2013  
L:34122 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2014  
L:34126 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2014  
L:34131 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2014  
L:34132 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2014  
L:34134 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2014  
L:34136 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2014  
L:34137 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2014  
L:34138 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2014  
L:34163 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2015  
L:34164 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2015  
L:34187 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2016  
L:34227 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2018  
L:34229 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2018  
L:34230 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2018  
L:34253 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2019  
L:34277 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2020  
L:34278 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2020  
L:34279 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2020  
L:34305 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2021  
L:34306 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2021

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L:34319 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2022

L:34346 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2023

L:34374 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2024